

of: htpan08xx.dat check: 8776 from: 1 to: 1643
symbols 1 to: 547.
of: Htpan08xx.Dat check: -1 from: 1 to: 1643 August 9, 1994 15:18
(entation)

!8.Gp_All check: 3291 from: 1 to: 279

MMU06948 1

✓ **Mus musculus Fas ligand mRNA, complete cds.** NCBI gi: 473565.
29-APR-1994

U06948

Mus musculus Eucaryotae; Metazoa; Chordata; Vertebrata;
Gnathostomata; Mammalia; Eutheria; Rodentia; Myomorpha; Muridae; . .

Symbol comparison table: Gencoredisk:[Gcgcore.Data.Rundata]Swgappep.Cmp
CompCheck: 1254

Gap Weight: 3.000 Average Match: 0.540
Length Weight: 0.100 Average Mismatch: -0.396

Quality: 113.1 **Length:** 300
Ratio: 0.427 **Gaps:** 7

Percent Similarity: 45.276 Percent Identity: 22.047

Htpan08xx.Pep x U06948.Gp_All August 9, 1994 15:29 ..

TRANSLATE of: htpan08xx.dat **check:** 8776 **from:** 1 **to:** 1643
generated symbols 1 to: 547.

**REFORMAT of: Htpan08xx.Dat check: -1 from: 1 to: 1643 August 9, 1994 15:18
(No documentation)**

to: U06948.Gp_All check: 3291 from: 1 to: 279

•LOCUS MMU06948_1
DEFINITION Mus musculus Fas ligand mRNA, complete cds. NCBI gi: 473565.
DATE 29-APR-1994
ACCESSION U06948
ORGANISM Mus musculus Eucaryotae; Metazoa; Chordata; Vertebrata;
Gnathostomata; Mammalia; Eutheria; Rodentia; Myomorpha; Muridae; .

Symbol comparison table: Gencoredisk:[Gcgcore.Data.Rundata]Swgappep.Cmp
CompCheck: 1254

Gap Weight: 3.000 Average Match: 0.540
Length Weight: 0.100 Average Mismatch: -0.396

Quality: 113.1 Length: 300
Ratio: 0.427 Gaps: 7
Percent Similarity: 45.276 Percent Identity: 22.047

Htpan08xx.Pep x U06948.Gp All August 9, 1994 15:29 ..

9	LQQSDSDRFMAMMEVQGGPSLG.....	QTCVLIVIFTVLLQSLCV	48
15	VDSSATSSWAPPGSVFP <small>CPCPSCGPRGRDORR</small> PPPPPPVSP LPPPSQPLPL		64
49	AVTYVYFTNELLKQM QDKYSKSGIACFLKEDDSYWD PNEESMN SPCWQVK		98
65	PP....LTPLKKKDHTNLVLPVVFFMVLV.....	ALVGMLGMYQLF	103
99	WQLRQLVRKMLRTSEETISTVQE KQQNISPLVRERGPQRVA AHITGTRG		148
104	HLQKELAELREFTNQSLKVSSFEKQIANPSTPSEKKEPRSVAHLTGNPHS		153
149	RSNTLSSPNSKNEKALGRKINSWE SSRS GHSFLSNLHLRN GELVIHEKG		198
154	RSIPLWEWT.....	YGTALISGVKYKKGGLVINETGL	186
199	YYIYSQTYFRFQEEIKENTKNDKQM VQYIMKYTS .YPDPI LLMKSARNSC		247
187	<u>YFVYSKVYFR</u>GQSCNNQPLNHKVYMRNSKYPEDLVLM ECKERLYN		230
248	WSKDAEYGLSIYQGGIFELKENDRIFVSVTNEHLIDMDHEASFFGAFLV		297
231	CTTGQIWA.HSSYLGAVFNLTSAHDLYVNISQLSLINFEESKTFFGLYKL		279